

## SEQUENCE LISTING

5 <110> Jackson, David C  
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 Walker, John  
 <120> T Helper Cell Epitopes  
 10 <130> 501902/MRO  
 <150> US 09/890650  
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 15 <150> PCT/AU00/00070  
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 <150> AU PP8533  
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 20 <150> AU PQ2013  
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&lt;400&gt; 21

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Thr

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 Ser Arg Ala Arg His Ser Thr Thr Ser Ala Gln Arg Ser Thr His Tyr  
 50 35 40 45  
 Asp Pro Arg Thr Ser Asp Arg Pro Val Ser Tyr Thr Met Asn Arg Thr  
 55 50 55 60  
 Arg Ser Arg Lys Gln Thr Ser His Arg Leu Lys Asn Ile Pro Val His  
 65 70 75 80  
 60 Gly Asn His Glu Ala Thr Ile Gln His Ile Pro Glu Ser Val Ser Lys  
 85 90 95  
 Gly Ala Arg Ser Gln Ile Glu Arg Arg Gln Pro Asn Ala Ile Asn Ser

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5	Gly Ser His Cys Thr Trp Leu Val 115	Leu Trp Cys Leu Gly Met Ala Ser 120	
10	Leu Phe Leu Cys Ser Lys Ala Gln Ile His Trp Asp Asn Leu Ser Thr 130		140
15	Ile Gly Ile Ile Gly Thr Asp Asn Val His Tyr Lys Ile Met Thr Arg 145	150	155 160
20	Pro Ser His Gln Tyr Leu Val Ile Lys Leu Ile Pro Asn Ala Ser Leu 165	170	175
25	Ile Glu Asn Cys Thr Lys Ala Glu Leu Gly Glu Tyr Glu Lys Leu Leu 180	185	190
30	Asn Ser Val Leu Glu Pro Ile Asn Gln Ala Leu Thr Leu Met Thr Lys 195	200	205
35	Asn Val Lys Pro Leu Gln Ser Leu Gly Ser Gly Arg Arg Gln Arg Arg 210	215	220
40	Phe Ala Gly Val Val Leu Ala Gly Val Ala Leu Gly Val Ala Thr Ala 225	230	235 240
45	Ala Gln Ile Thr Ala Gly Ile Ala Leu His Gln Ser Asn Leu Asn Ala 245	250	255
50	Gln Ala Ile Gln Ser Leu Arg Thr Ser Leu Glu Gln Ser Asn Lys Ala 260	265	270
55	Ile Glu Glu Ile Arg Glu Ala Thr Gln Glu Thr Val Ile Ala Val Gln 275	280	285
60	Gly Val Gln Asp Tyr Val Asn Asn Glu Leu Val Pro Ala Met Gln His 290	295	300
65	Met Ser Cys Glu Leu Val Gly Gln Arg Leu Gly Leu Arg Leu Leu Arg 305	310	315 320
70	Tyr Tyr Thr Glu Leu Leu Ser Ile Phe Gly Pro Ser Leu Arg Asp Pro 325	330	335
75	Ile Ser Ala Glu Ile Ser Ile Gln Ala Leu Ile Tyr Ala Leu Gly Gly 340	345	350
80	Glu Ile His Lys Ile Leu Glu Lys Leu Gly Tyr Ser Gly Ser Asp Met		



	355	360	365
5	Ile Ala Ile Leu Glu Ser Arg Gly Ile Lys Thr Lys Ile Thr His Val 370 375 380		
10	Asp Leu Pro Gly Lys Phe Ile Ile Leu Ser Ile Ser Tyr Pro Thr Leu 385 390 395 400		
15	Ser Glu Val Lys Gly Val Ile Val His Arg Leu Glu Ala Val Ser Tyr 405 410 415		
20	Asn Ile Gly Ser Gln Glu Trp Tyr Thr Thr Val Pro Arg Tyr Ile Ala 420 425 430		
25	Thr Asn Gly Tyr Leu Ile Ser Asn Phe Asp Glu Ser Ser Cys Val Phe 435 440 445		
30	Val Ser Glu Ser Ala Ile Cys Ser Gln Asn Ser Leu Tyr Pro Met Ser 450 455 460		
35	Pro Leu Leu Gln Gln Cys Ile Arg Gly Asp Thr Ser Ser Cys Ala Arg 465 470 475 480		
40	Thr Leu Val Ser Gly Thr Met Gly Asn Lys Phe Ile Leu Ser Lys Gly 485 490 495		
45	Asn Ile Val Ala Asn Cys Ala Ser Ile Leu Cys Lys Cys Tyr Ser Thr 500 505 510		
50	Ser Thr Ile Ile Asn Gln Ser Pro Asp Lys Leu Leu Thr Phe Ile Ala 515 520 525		
55	Ser Asp Thr Cys Pro Leu Val Glu Ile Asp Gly Ala Thr Ile Gln Val 530 535 540		
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	Asn Ala Leu Lys Lys Leu Asp Asp Ala Lys Val Leu Ile Asp Ser Ser 580 585 590		
	Asn Gln Ile Leu Glu Thr Val Arg Arg Ser Ser Phe Asn Phe Gly Ser 595 600 605		
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	610	615	620
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